

=> d his full

(FILE 'HOME' ENTERED AT 09:55:34 ON 15 OCT 2008)

FILE 'REGISTRY' ENTERED AT 09:55:47 ON 15 OCT 2008

L1 STRUCTURE UPLOADED  
D  
L2 1 SEA SSS SAM L1  
L3 55 SEA SSS FUL L1

FILE 'CAPLUS' ENTERED AT 09:57:25 ON 15 OCT 2008

L4 19 SEA ABB=ON PLU=ON L3  
L5 12 SEA ABB=ON PLU=ON L4 AND GLYCOPHINGOLIPIDS  
D BIB ABS HITSTR L5 1-12

FILE 'STNGUIDE' ENTERED AT 09:58:36 ON 15 OCT 2008

FILE 'CAPLUS' ENTERED AT 10:06:05 ON 15 OCT 2008

L6 1 SEA ABB=ON PLU=ON L4 AND VIRUS  
D SCAN  
D HSI

FILE 'STNGUIDE' ENTERED AT 10:08:52 ON 15 OCT 2008

FILE 'CAPLUS' ENTERED AT 10:11:58 ON 15 OCT 2008

D L6 HITSTR  
L7 0 SEA ABB=ON PLU=ON L4 AND VIRAL  
L8 4850 SEA ABB=ON PLU=ON GLYCOPHINGOLIPIDS  
L9 195 SEA ABB=ON PLU=ON L8 AND VIRUS  
L10 46 SEA ABB=ON PLU=ON L8 AND ANTIVIRAL  
E TEYTON L/AU  
L11 112 SEA ABB=ON PLU=ON "TEYTON L"/AU OR "TEYTON LUC"/AU  
L12 19 SEA ABB=ON PLU=ON L11 AND GLYCOLIPID  
L13 1 SEA ABB=ON PLU=ON L12 AND VIRUS  
D SCAN  
L14 10 SEA ABB=ON PLU=ON L11 AND VIRUS  
D SCAN

10553695

=> d his full

(FILE 'HOME' ENTERED AT 15:10:09 ON 15 OCT 2008)

FILE 'REGISTRY' ENTERED AT 15:10:20 ON 15 OCT 2008

L1 STRUCTURE UPLOADED

D

L2 55 SEA SSS FUL L1

FILE 'CAPLUS' ENTERED AT 15:10:55 ON 15 OCT 2008

L3 19 SEA ABB=ON PLU=ON L2

L4 1 SEA ABB=ON PLU=ON L3 AND VIRUS

D SCAN

L5 1849 SEA ABB=ON PLU=ON SPHINGOMONAS

L6 13 SEA ABB=ON PLU=ON L5 AND VIRUS

L7 26 SEA ABB=ON PLU=ON L5 AND GLYCOLIPIDS

L8 2 SEA ABB=ON PLU=ON L7 AND VIRUS

D SCAN

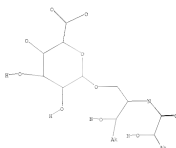
D BIB ABS 1-2

L9 0 SEA ABB=ON PLU=ON SPHINGOMONAS GSL-1

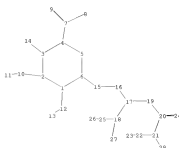
L10 21 SEA ABB=ON PLU=ON GSL-1

L11 1 SEA ABB=ON PLU=ON L10 AND VIRUS

D SCAN



C



29

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chain nodes :
  7  8  9  10  11  12  13  14  15  16  17  18  19  20  21  22  23  24  25  26  27  28  29
ring nodes :
  1  2  3  4  5  6
chain bonds :
  1-12  2-10  3-14  4-7  6-15  7-8  7-9  10-11  12-13  15-16  16-17  17-18  17-19  18-25  18-27
  19-20  20-21  20-24  21-22  21-28  22-23  25-26
ring bonds :
  1-2  1-6  2-3  3-4  4-5  5-6
exact/norm bonds :
  1-2  1-6  1-12  2-3  2-10  3-4  3-14  4-5  5-6  6-15  7-8  7-9  15-16  17-19  18-25  18-27
  19-20  20-24  21-22  21-28
exact bonds :
  4-7  10-11  12-13  16-17  17-18  20-21  22-23  25-26

Match level :
1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 10:CLASS 11:CLASS
12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 17:CLASS 18:CLASS 19:CLASS 20:CLASS
21:CLASS 22:CLASS 23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS 29:CLASS

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